

ABSTRACT

It is intended to provide a means of predicting genetic risk for hypertension at a high accuracy and high prediction possibility. Namely, risk for hypertension is
5 predicted by a method involving the following steps: (i) the step of analyzing two or more polymorphisms selected from among 4 gene polymorphisms having been revealed as relating to hypertension; (ii) the step of determining the genotype of a nucleic acid sample based on the polymorphism data obtained in the above step; and
10 (iii) the step of predicting the genetic risk for hypertension from the genotype thus determined.